BIO. CHNOLOGY
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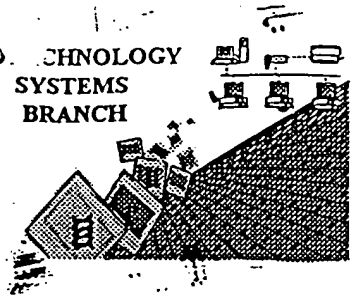

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RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/028,952
Source: OIPE
Date Processed by STIC: 1-23-03

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FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
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Revised 01/29/2002

Raw Sequence Listing Error Summary

ERROR DETECTED **SUGGESTED CORRECTION** SERIAL NUMBER: 10/028,952

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1	Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor **after** creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2	Invalid Line Length	The rules require that a line **not exceed** 72 characters in length. This includes white spaces.
3	Misaligned Amino Numbering	The numbering under each 5th amino acid is misaligned. Do **not** use tab codes between numbers; use **space characters**, instead.
4	Non-ASCII	The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. **Please ensure your subsequent submission is saved in ASCII text.**
5	Variable Length	Sequence(s)____contain n's or.Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6	PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**
7	Skipped Sequences (OLD RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for **each** skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.
8	Skipped Sequences (NEW RULES)	Sequence(s) _____ missing. If **intentional**, please insert the following lines for **each** skipped sequence. <210> sequence id number <400> sequence id number 000
9	Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of **n** or **Xaa**, and which residue **n** or **Xaa** represents.
10	Invalid <213> Response	Per 1.823 of Sequence Rules, the only **valid** <213> responses are: Unknown, Artificial *anything*, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or contains Artificial.
11	Use of <220>	Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response contains the word "Artificial" or "Unknown." Please explain source of genetic material in <220> to <223> section, i.e., why you chose Artificial or Unknown. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
12	PatentIn "bug"	Please do not use "Copy to Disk" function of PatentIn. In PatentIn 2.x it causes a corrupted file and in PatentIn 3.x you may lose your hard returns in the sequence listing. Instead, please use "Windows Explorer" or any other manual means to copy file to floppy disk.

Does Not Comply
Corrected Diskette Needed
See Additional pages 1-13

OIPE

RAW SEQUENCE LISTING DATE: 01/23/2003
PATENT APPLICATION: **US/10/028,952** TIME: 13:55:27

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Output Set: **N:\CRF4\01232003\J028952.raw**

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      7 <130> FILE REFERENCE: 4273.3USW1
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PATENT APPLICATION: **US/10/028,952** TIME: 13:55:27

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202 agacagcttg gctcttgccc ctgacaggat actgagccaa gccctgcctg tggccaagcc 1860
203 ctgagtggcc actgccaagc tgcggggaag ggtcctgagc aggggcatct gggaggctct 1920
204 ggctgccttc tgcatttatt tgccttttтt ctttttctct tgcttctaag gggtggtggc 1980
205 caccactgtt tagaatgacc cttgggaaca gtgaacgtag agaattgttt ttagcagagt 2040
206 ttgtgaccaa agtcagagtg gatcatggtg gtttggcagc agggaatttg tcttgttgga 2100
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208 gatgcagatg gccaagctcc caccctgggt attcaaaaac ggcagacaca acatgttcct 2220
209 ccacgcggct cactcgatgc ctgcaggccc cagtgtgtgc ctcaactgat tctgacttca 2280
210 ggaaaagtaa aaaaaaaaaa aaaaaactcg agaagctttg gacttcttcg cca        2333
```

RAW SEQUENCE LISTING DATE: 01/23/2003
PATENT APPLICATION: **US/10/028,952** TIME: 13:55:27

Input Set : **A:\4273-3u1.app**
Output Set: **N:\CRF4\01232003\J028952.raw**

<210> SEQ ID NO: 5
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<212> TYPE: DNA
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 01/23/2003
PATENT APPLICATION: **US/10/028,952** TIME: 13:55:28

Input Set : **A:\4273-3u1.app**
Output Set: **N:\CRF4\01232003\J028952.raw**

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:9; Xaa Pos. 16,41,44,48,64,96,112,118,131,144,160,168,173,192,198,208
Seq#:9; Xaa Pos. 227,240,256,264,276,288,302,304,336,352,383,399,431,447
Seq#:9; Xaa Pos. 479,495,520,522,527,543,544,552,556,558,575,591,602,622
Seq#:9; Xaa Pos. 623,639,671,687,719,735
Seq#:10; Xaa Pos. 10,13,32,48,80,87,96,100,128,137,142,144,166,176,192,196
Seq#:10; Xaa Pos. 224,233,240,245,270,272,288,320,336,352,353,363,367,368
Seq#:10; Xaa Pos. 371,383,384,385,410,416,432,438,464,480,487,495,496,497
Seq#:10; Xaa Pos. 506,512,513,522,528,550,554,560,574,576,589,591,608,624
Seq#:10; Xaa Pos. 652,656,657,662,663,672,702,704,718,720,723,734,743,752

VERIFICATION SUMMARY DATE: 01/23/2003
PATENT APPLICATION: **US/10/028,952** TIME: 13:55:28

Input Set : **A:\4273-3u1.app**
Output Set: **N:\CRF4\01232003\J028952.raw**

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:500 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ ID#:9
L:500 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:506 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:32
L:509 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:48
L:515 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:80
L:518 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:96
L:521 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:112
L:524 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:128
L:527 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:144
L:530 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:160
L:533 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:176
L:536 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:192
L:542 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:224
L:545 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:240
L:548 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:256
L:551 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:272
L:554 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:288
L:560 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:320
L:563 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:336
L:569 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:368
L:572 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:384
L:578 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:416
L:581 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:432
L:587 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:464
L:590 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:480
L:596 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:512
L:599 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:528
L:602 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:544
L:605 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:560
L:608 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:576
L:611 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:592
L:614 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:608
L:617 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:624
L:623 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:656
L:626 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:672
L:632 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:704
L:635 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:720
L:899 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ ID#:10
L:899 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:902 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:16
L:905 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:32
L:911 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:64
L:914 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:80
L:917 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:96
L:920 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:112
L:923 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:128
L:929 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:160

VERIFICATION SUMMARY DATE: 01/23/2003
PATENT APPLICATION: **US/10/028,952** TIME: 13:55:28

Input Set : **A:\4273-3u1.app**
Output Set: **N:\CRF4\01232003\J028952.raw**

L:932 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:176
L:935 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:192
L:938 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:208
L:941 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:224
L:944 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:240

Additional page 1

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See item # 9 on Error Summary Sheet. Each "Xaa" requires feature containing numeric identifiers <220>, <221>, <222>, and <223>.

If all Xaa's in the sequence are equal to the same thing. One feature with a range of the whole sequence in numeric identifier <222> is all that is necessary.

Example:

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            20                  25                  30

Leu Val Trp His Gly Glu Glu Thr Xaa Glu Val Xaa Asn Lys Trp Xaa
        35                  40                  45

Ala Pro Gly Ala Pro Pro Val Ser Pro Arg Gly Ala Arg Gly Gly Xaa
    50                  55                  60

Arg Pro Cys Gly Pro Pro Val Lys Tyr His Tyr Ser Asp Arg Phe Thr
65                  70                  75                  80

Asp Pro Val Arg Arg Gly Gly Glu Pro Arg Gly Ala Leu Ala Ser Xaa
                85                  90                  95

Ala Lys Arg Pro Ala Ala Arg Arg Pro Gly Ala Thr Arg Ser Gly Xaa
            100                 105                 110

Ala Arg Trp Gly Val Xaa Leu Gly Arg Tyr Thr Cys Gln Thr Val Gln
        115                 120                 125

Val Ser Xaa Gly Glu Leu Arg Glu Asp Arg Asn Leu Pro Trp Ser Xaa
    130                 135                 140

Arg Ala Lys Ala Arg Leu Ile Leu Ile Phe Ser Thr Asn Thr Asp Xaa
145                 150                 155                 160

Ser Gly Ala Ser Arg Ser Phe Xaa Pro Phe Gly Phe Xaa Ala Gly Val
                165                 170                 175

Arg Lys Val Thr Thr Gly Ile Thr Gly Leu Trp Arg Pro Ser Val Xaa
            180                 185                 190

Ser Asp Val Ala Phe Xaa Ser Phe Asp Val Gly Ser Ser Tyr His Xaa
        195                 200                 205

Ala Glu Phe Thr Lys Arg Trp Ile Val His Pro Leu Ile Gly Asn Ser
    210                 215                 220

Trp Asp Xaa Thr Val Val Arg Gln Val Ser Phe Thr Leu Leu Met Xaa
225                 230                 235                 240

Cys Cys Cys His Gly Asn Pro Ala Gln Tyr Glu Arg Asn Arg Arg Xaa
```

245 250 255

His Leu Val Tyr Val Leu Gly Xaa Gly Ala Asn Gly Ala Lys Leu Ser
260 265 270

Val Gly Leu Xaa Leu Asn Ala Ser Lys Ser Glu Ser Arg Pro Gly Xaa
275 280 285

Thr Ile Arg Gln Arg Arg Gly Ala Ser Val Gly Leu Gly Xaa Pro Xaa
290 295 300

Arg Leu Ser Pro Pro Ala Gly Arg Pro Pro Leu His Ala Pro Arg Arg
305 310 315 320

Gly Arg Ala Arg Ala Pro Pro Arg Ala Gly Thr Gly Val Arg Cys Xaa
325 330 335

Val Pro Phe Val Leu Gly Asn Gly Ala Arg Pro Glu Arg Arg Pro Xaa
340 345 350

Arg Pro Ser Arg Thr Ala Arg Ser Trp Gly Thr Trp Arg Thr Ser Ile
355 360 365

Ser Ser Pro Gln Pro Gly Lys Leu Arg Ser Pro Phe Leu Gln Xaa Gln
370 375 380

Leu Thr Gln Pro Glu Thr His Phe Gly Arg Glu Pro Ala Ala Xaa Ser
385 390 395 400

Arg Pro Arg Ala Asp Leu Pro Ala Glu Glu Pro Ala Pro Ser Pro Pro
405 410 415

Cys Leu Val Gln Ala Glu Glu Glu Ala Val Tyr Glu Glu Pro Xaa Glu
420 425 430

Gln Glu Thr Phe Tyr Glu Gln Pro Pro Leu Val Gln Gln Gln Xaa Gly
435 440 445

Ser Glu His Ile Asp His His Ile Gln Gly Gln Gly Leu Ser Gln Gly
450 455 460

Leu Cys Ala Arg Ala Leu Tyr Asp Tyr Gln Ala Ala Asp Asp Xaa Glu
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Ile Ser Phe Asp Pro Glu Asn Leu Ile Thr Gly Ile Glu Val Xaa Glu
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Gly Trp Trp Arg Gly Tyr Gly Pro Asp Gly His Phe Gly Met Pro Ala
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Asn Tyr Val Glu Leu Ile Glu Xaa Gly Xaa Gly His Ile Leu Xaa Phe
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Pro Ser Gln Thr Trp Leu Pro Tyr Cys Trp Lys Arg Arg Pro Xaa Xaa
530 535 540

His Ser Ala Leu Phe Gln Glu Xaa Asp Pro Gln Xaa Gly Xaa Leu Arg

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545                 550                 555                 560

Ala Pro Ser Gly Leu Ala Asp Ser Ala Cys His Pro Lys Cys Xaa Asn
                565                 570                 575

Gly Leu Val Ile Pro Thr His Pro Ser Cys Ile Pro Arg Pro Xaa Thr
            580                 585                 590

Ala Trp Leu Leu Pro Leu Thr Gly Tyr Xaa Ala Lys Pro Cys Trp Pro
        595                 600                 605

Ser Pro Glu Trp Pro Leu Pro Ser Cys Gly Glu Gly Ser Xaa Xaa Gly
    610                 615                 620

Ala Ser Gly Arg Leu Trp Leu Pro Ser Ala Phe Ile Cys Leu Xaa Phe
625                 630                 635                 640

Ser Leu Ala Ser Lys Gly Trp Trp Pro Pro Leu Phe Arg Met Leu Gly
                645                 650                 655

Asn Ser Glu Arg Arg Glu Leu Phe Leu Ala Glu Phe Val Thr Xaa Val
            660                 665                 670

Arg Val Asp His Gly Gly Leu Ala Ala Gly Asn Leu Ser Cys Xaa Leu
        675                 680                 685

Leu Cys Ala Pro His Ser Ile Ser Leu Ser Leu Cys Leu Gly Gly Lys
    690                 695                 700

Trp Gly Cys Arg Trp Pro Ser Ser His Pro Gly Tyr Ser Lys Xaa Ala
705                 710                 715                 720

Asp Thr Thr Cys Ser Ser Thr Arg Leu Thr Arg Cys Leu Gln Xaa Val
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Cys Ala Ser Thr Asp Ser Asp Phe Arg Lys Ser Lys Lys Lys Lys Lys
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Ile Arg His Glu His Gly Glu Glu Thr Xaa Glu Val Xaa Asn Lys Glu
  1               5                  10                  15

Ala Pro Gly Ala Pro Pro Val Ser Pro Arg Gly Ala Arg Gly Gly Xaa
             20                  25                  30

Arg Arg Pro Cys Gly Pro Pro Val Lys Tyr His Tyr Ser Asp Arg Xaa
         35                  40                  45

Thr Asp Pro Val Arg Arg Gly Gly Glu Pro Arg Gly Ala Leu Ala Gly
     50                  55                  60

Ala Lys Arg Pro Ala Ala Arg Arg Pro Gly Ala Thr Arg Ser Gly Xaa
 65                  70                  75                  80

Ser Ala Arg Trp Gly Val Xaa Leu Gly Arg Tyr Thr Cys Gln Thr Xaa
                 85                  90                  95

Gln Val Ser Xaa Gly Glu Leu Arg Glu Asp Arg Asn Leu Pro Trp Arg
            100                 105                 110

Arg Ala Lys Ala Arg Leu Ile Leu Ile Phe Ser Thr Asn Thr Asp Xaa
        115                 120                 125

Glu Ser Gly Ala Ser Arg Ser Phe Xaa Pro Phe Gly Phe Xaa Ala Xaa
    130                 135                 140

Val Arg Lys Val Thr Thr Gly Ile Thr Gly Leu Trp Arg Pro Ser His
145                 150                 155                 160

Ser Asp Val Ala Phe Xaa Ser Phe Asp Val Gly Ser Ser Tyr His Xaa
                165                 170                 175

Glu Ala Glu Phe Thr Lys Arg Trp Ile Val His Pro Leu Ile Gly Xaa
            180                 185                 190

Ser Trp Val Xaa Thr Val Val Arg Gln Val Ser Phe Thr Leu Leu Met
        195                 200                 205

Cys Cys Cys His Gly Asn Pro Ala Gln Tyr Glu Arg Asn Arg Arg Xaa
    210                 215                 220

Arg His Leu Val Tyr Val Leu Gly Xaa Gly Ala Asn Gly Ala Lys Xaa
225                 230                 235                 240

Ser Val Gly Leu Xaa Leu Asn Ala Ser Lys Ser Glu Ser Arg Pro Gly
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245 250 255

Thr Ile Arg Gln Arg Arg Gly Ala Ser Val Gly Leu Gly Xaa Pro Xaa
260 265 270

Pro Arg Leu Ser Pro Pro Ala Gly Arg Pro Pro Pro Ser Thr Arg Xaa
275 280 285

Arg Ala Gly Gly Arg Val Pro Arg Arg Ala Pro Gly Pro Gly Ser Ala
290 295 300

Glu Cys Pro Ser Ser Trp Glu Thr Gly Arg Gly Arg Lys Gly Gly Xaa
305 310 315 320

Pro Leu Ala Arg His Ala Pro His Val Arg Ala Arg Ala Glu Phe Xaa
325 330 335

Ser Ser Thr Ile His Asn Arg His Thr Ser Ala Cys Ile Phe Met Xaa
340 345 350

Xaa Ile Leu Phe Leu Trp Val Asp Ile Gln Xaa Trp Asp Cys Xaa Xaa
355 360 365

Thr Trp Xaa Phe Tyr Phe Trp Phe Ile Glu Lys Ser Ser Tyr Xaa Xaa
370 375 380

Xaa Arg Leu Tyr Lys Phe Thr Ser Leu Pro Ser Asp Phe Phe Lys Glu
385 390 395 400

Arg Met Val Trp Arg Asn Ala Pro His Xaa Tyr Pro Pro Phe Thr Xaa
405 410 415

Leu Leu Gln Asn Asp Phe Lys Gly Tyr Arg Tyr Leu Gln Val Ser Xaa
420 425 430

Arg Gln Ile Glu Tyr Xaa Asn Phe Cys Ile Arg Gly Thr Asp Phe Ile
435 440 445

Gln Ser Cys Met Asn Lys Asp Lys Cys Ser Arg Asp Leu Gln Ser Xaa
450 455 460

Asn Trp Lys Ser Gln Met Lys Tyr Ile Ser Ser Ser Thr Thr Ser Xaa
465 470 475 480

Ser Thr Glu Leu Ala Leu Xaa Ser Ser Leu Ile Pro Thr Tyr Xaa Xaa
485 490 495

Xaa Lys Gly Phe Ile Ser Asn Ile Leu Xaa Gly Ile Lys Ile Lys Xaa
500 505 510

Xaa Val Lys Leu Phe Ser Leu Ala Phe Xaa Phe Gln Asn Ile Lys Xaa
515 520 525

Pro Ser Ile Gly His Leu Tyr Cys Thr Arg His Cys Val Cys His Ser
530 535 540

Lys Met Phe Ser Trp Xaa Cys Ser Gln Xaa Phe Cys Arg Val Arg Xaa

545 550 555 560

Ser Leu Thr Val Val Arg Leu Phe Ser Lys Arg Asn Leu Xaa Thr Xaa
565 570 575

Phe Asn Leu Arg Lys Val Ser Asn Arg Thr Arg Thr Xaa Thr Xaa Ile
580 585 590

Thr Leu Gln Ile Ser Pro Tyr His Thr Ala Ser Thr Cys Ala Cys Xaa
595 600 605

Leu Ile Pro Gly Ser Cys Tyr Phe Pro Phe Tyr Phe Leu Ser Leu Xaa
610 615 620

Thr Thr Pro Phe Ser Pro His Phe Phe Ser Phe Phe Leu Ile Val Tyr
625 630 635 640

Ile Thr Asn Thr Cys Leu Ser Glu Gln Leu Ile Xaa His Lys Arg Xaa
645 650 655

Xaa Ser Thr Gly Glu Xaa Xaa Leu Ile Pro Val Ile Leu Ala Leu Xaa
660 665 670

Ala Lys Ala Gly Arg Ser Leu Glu Ser Arg Val Arg Asp Gln Pro Gln
675 680 685

His Gly Glu Thr Leu Ser Leu Gln Lys Asn Thr Lys Ile Xaa Pro Xaa
690 695 700

Val Leu Ala His Thr Cys Ser Leu Ser Tyr Ser Glu Gly Xaa Gly Xaa
705 710 715 720

Ile Asp Xaa Ala Gln Glu Val Glu Ala Ala Ala Val Arg Xaa Asp Ala
725 730 735

Ile Ala Leu Gln Pro Gly Xaa Glu Arg Glu Thr Leu Ser Gln Lys Xaa
740 745 750

Lys